

FastQC Report

Mon 14 Apr 2014

2831_4902_8529_H8PNLADXX_GS061B_R1.fq

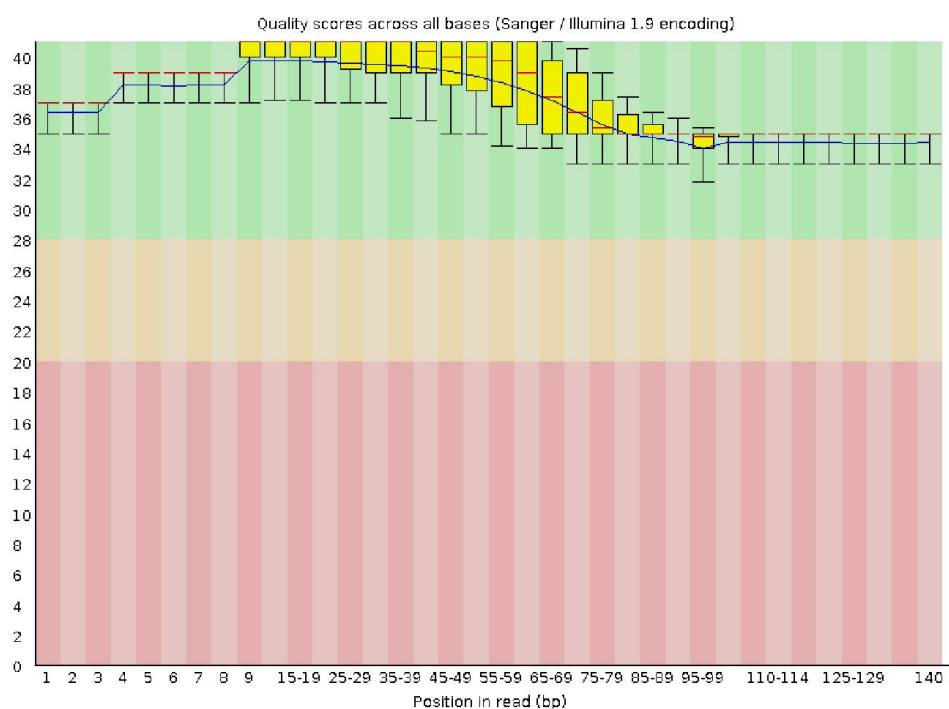
Summary

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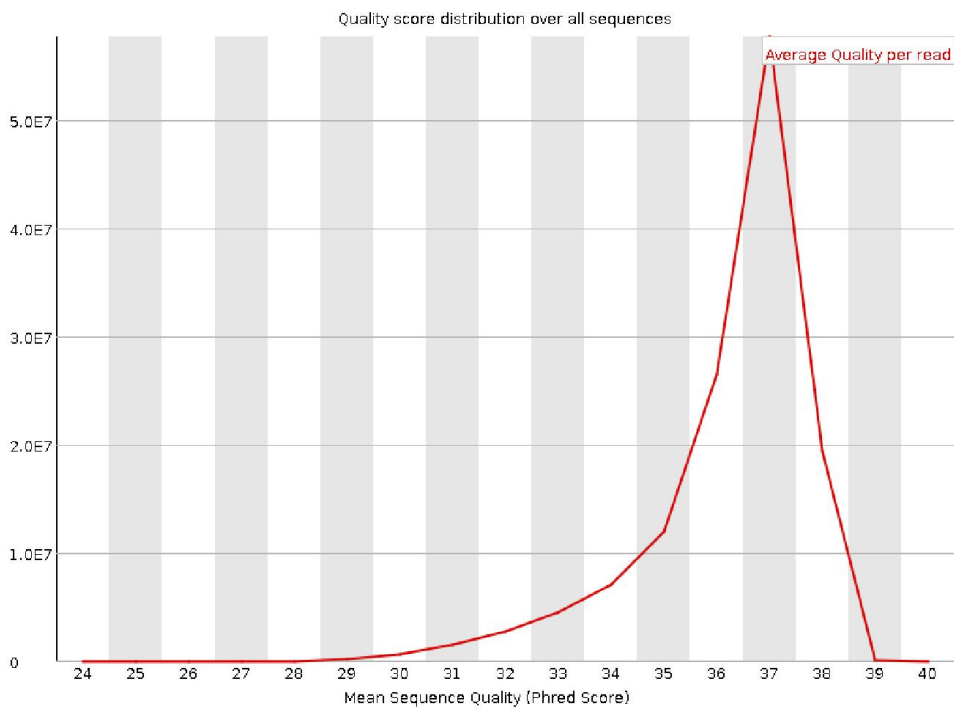
Basic Statistics

Measure	Value
Filename	2831_4902_8529_H8PNLADXX_GS061B_R1.fq
File type	Conventional base calls
Encoding	Sanger / Illumina 1.9
Total Sequences	133065110
Filtered Sequences	0
Sequence length	60-140
%GC	42

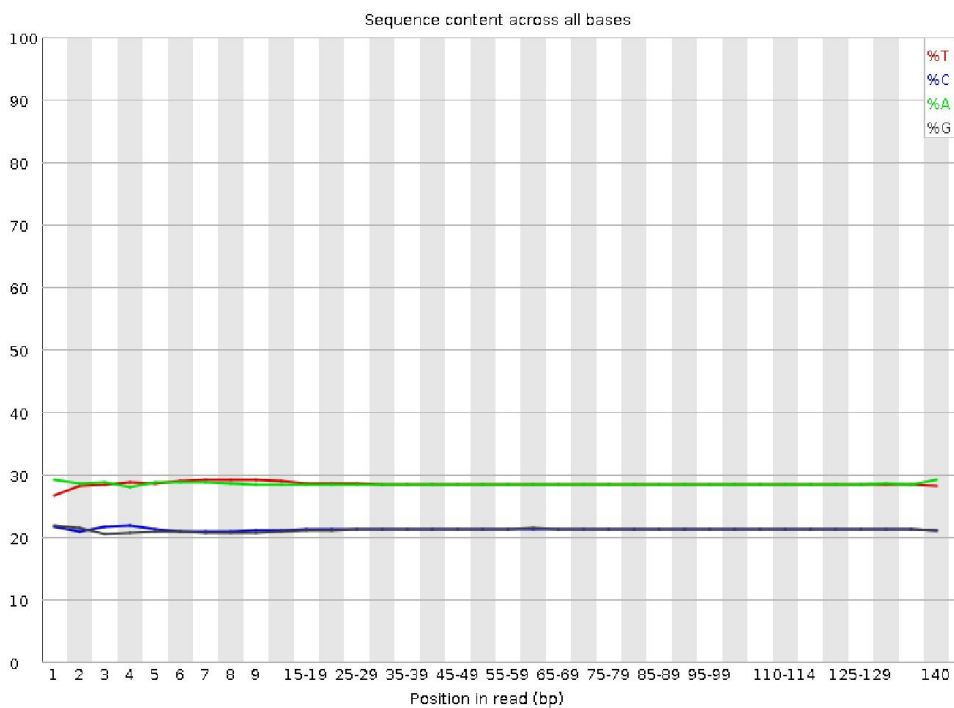
Per base sequence quality



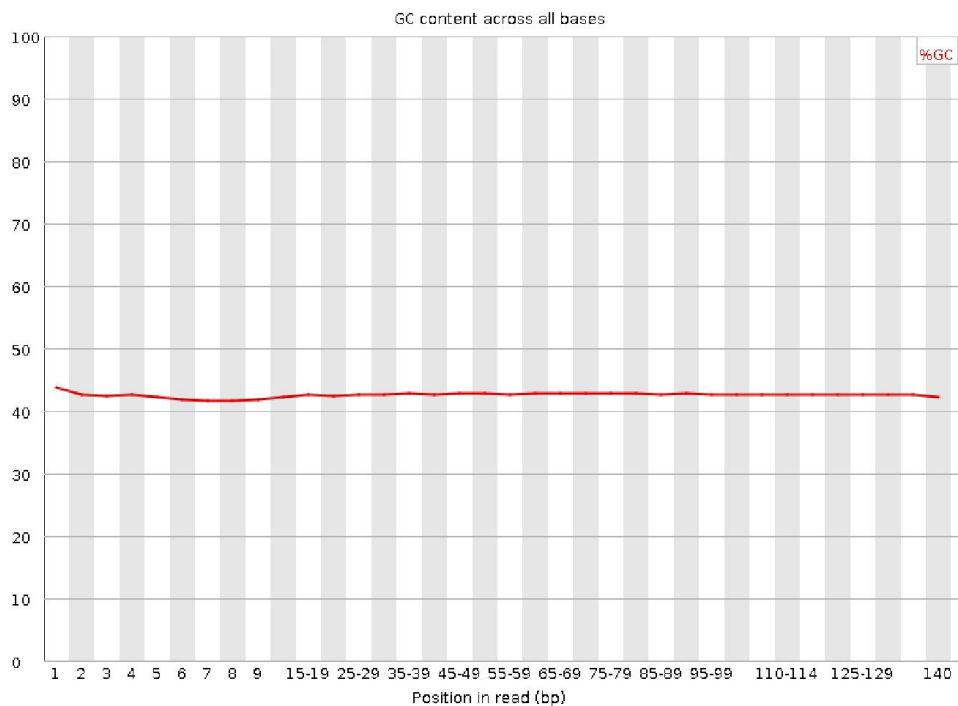
✓ Per sequence quality scores



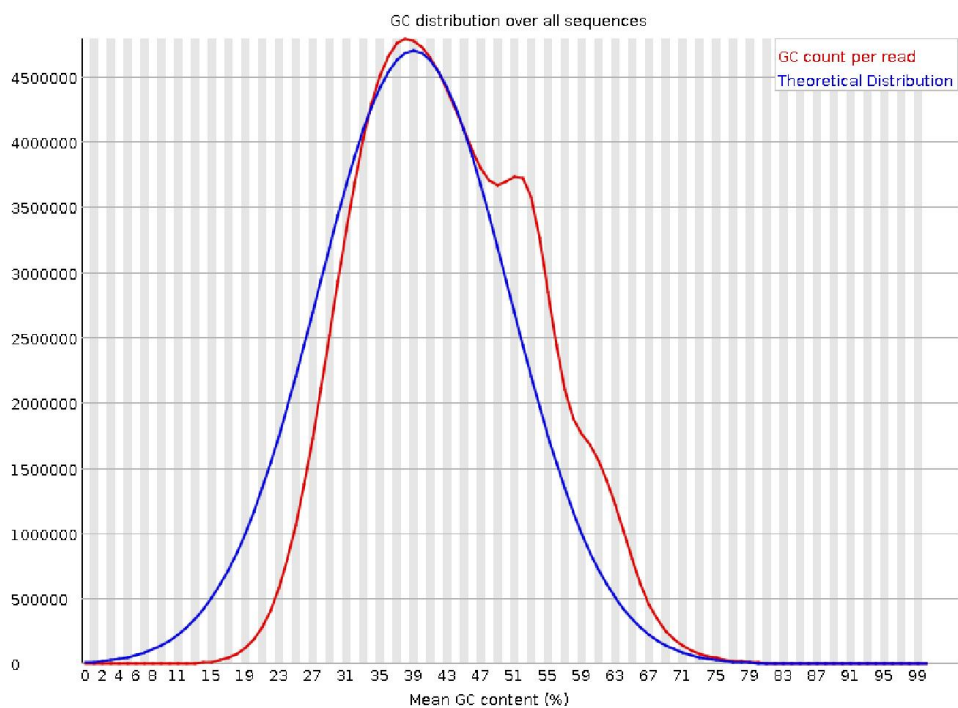
✓ Per base sequence content



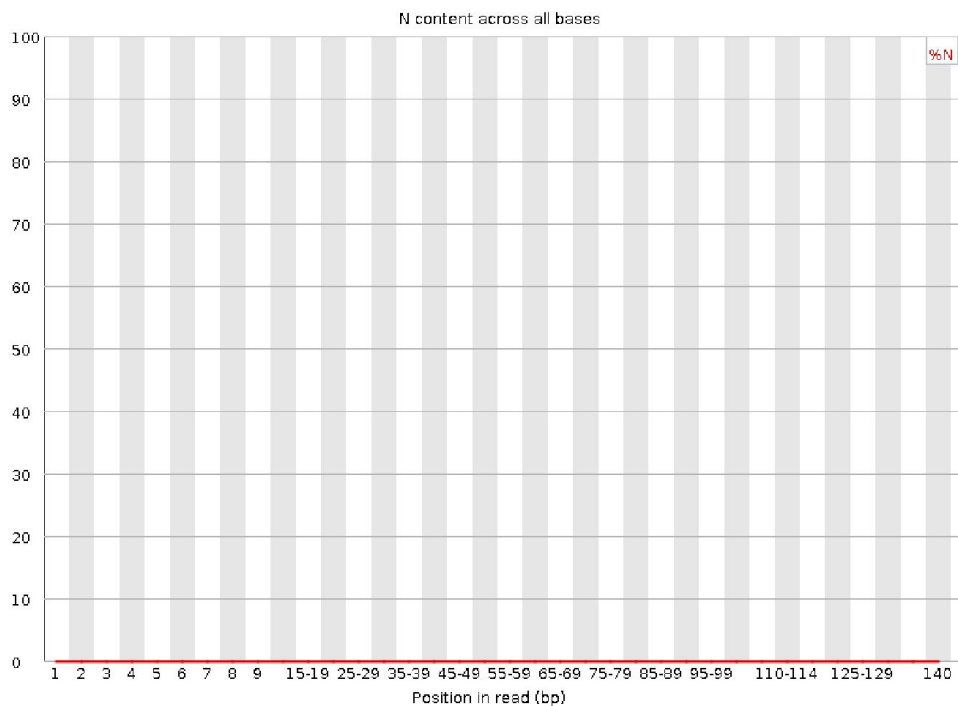
✓ Per base GC content



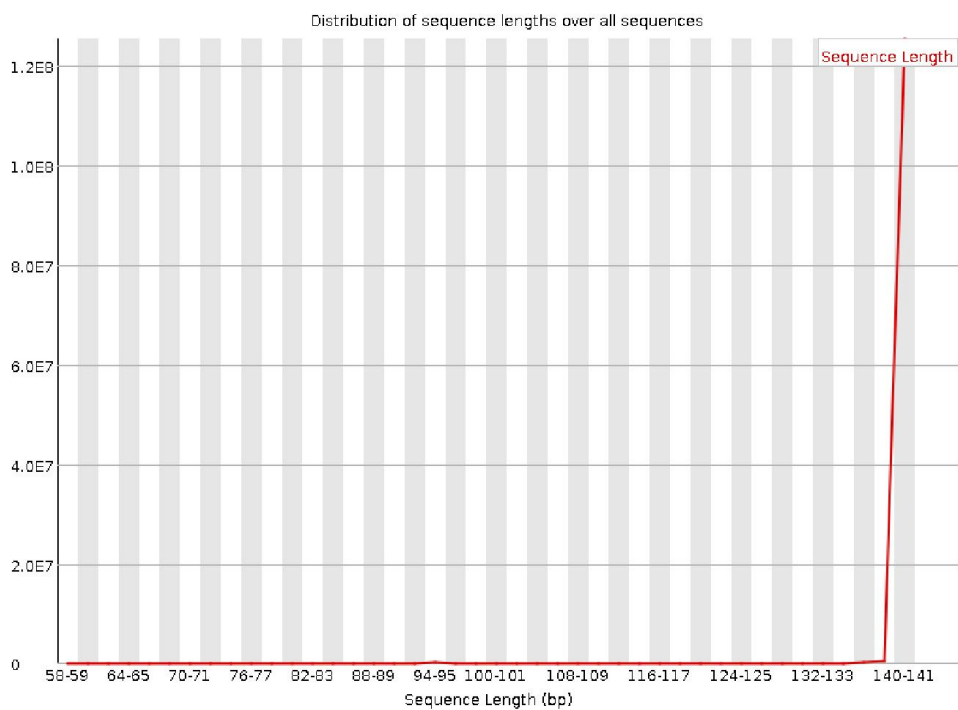
⚠ Per sequence GC content



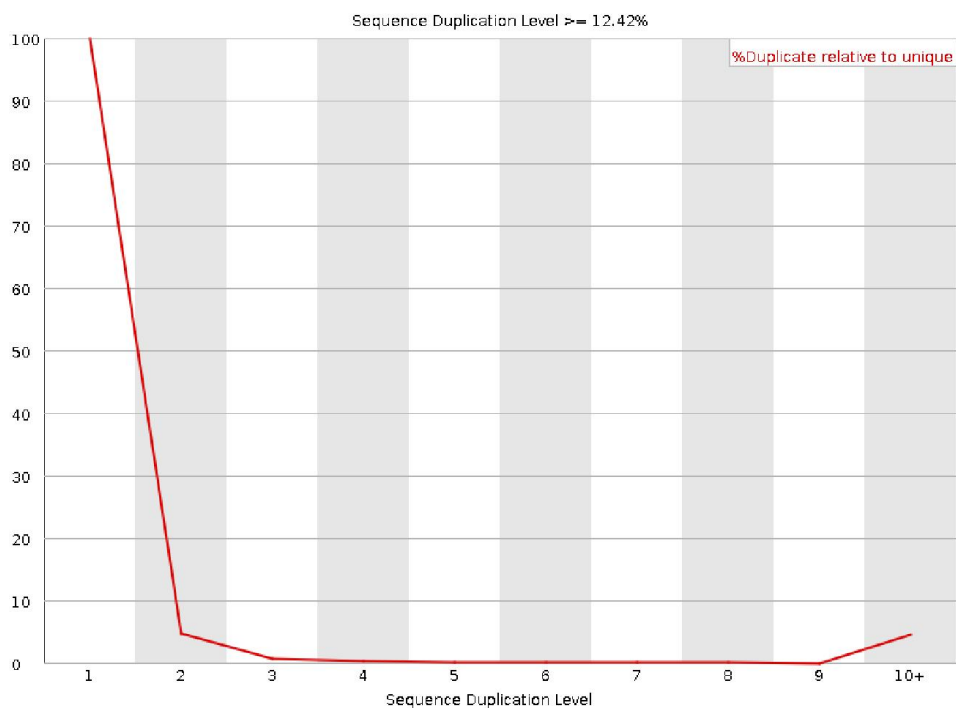
✅ Per base N content



Sequence Length Distribution



Sequence Duplication Levels



Overrepresented sequences

No overrepresented sequences



Kmer Content

No overrepresented Kmers

Produced by [FastQC](#) (version 0.10.1)